

Ad2	GNLTSQNVTTVTQPLKKTKS	SEQ ID NO: 42
Ad5	GNLTSQNVTTVSPPLKKTKS	SEQ ID NO: 43
Ad37	GSLTVNPK-----APLQVNTD	SEQ ID NO: 58
Ad8	GKLLTVNTE-----PPLHLTNN	SEQ ID NO: 66
Ad9	GKLLTVNAD-----PPLQLTNN	SEQ ID NO: 67
Ad15	GNLTVNTE-----PPLQLTNN	SEQ ID NO: 68

Alignment of the Corresponding Third  
Repeats of Ad Fiber Shaft Proteins

Figure 1A

Ad2	GAMITKLGAGLSFDNS	SEQ ID NO: 46
Ad5	KAMVPKLGTLGSLFDST	SEQ ID NO: 47
Ad37	IGINVRAREGLTFDND	SEQ ID NO: 48
Ad8	--VRVGEKGGLSFNDN	SEQ ID NO: 59
Ad9	--VRVGEKGGLSFNND	SEQ ID NO: 60
Ad15	--VRVGEKGGLSFNEA	SEQ ID NO: 61
Consensus	KLGXGLXFD	
	N	

Alignment of the Corresponding Last Full  
Repeats of Ad Fiber Shaft Proteins

Figure 1B

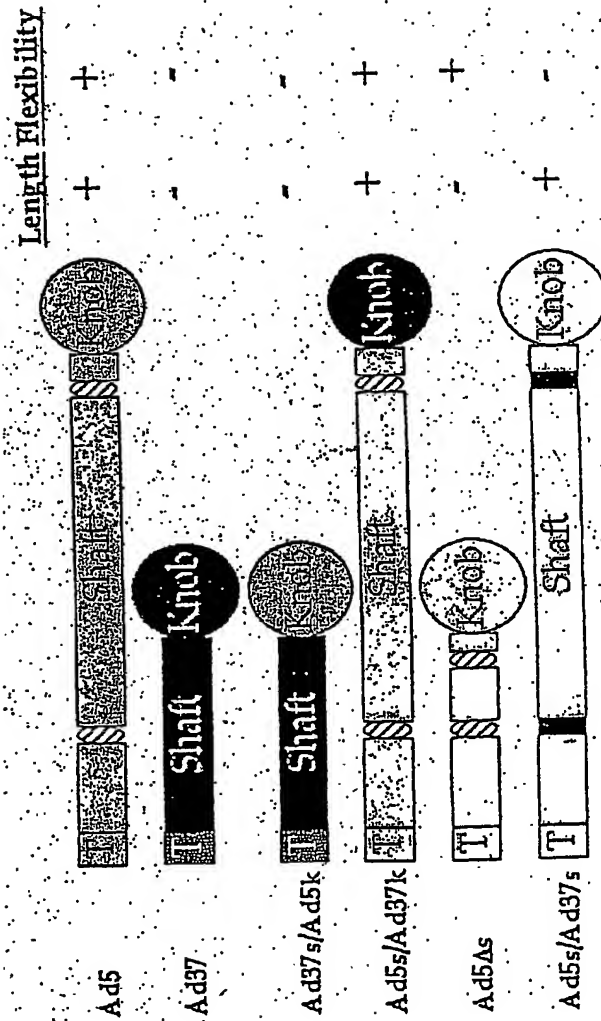


FIGURE 2